

SEQUENCE LISTING

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SLOOTS, THEO PEITER

<120> NEISSERIA GONORRHOEAE DETECTION

<130> 28472-0003

<140> 10/599,699

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<150> PCT/AU05/000500

<151> 2005-04-06

<150> AU 2004901890

<151> 2004-04-08

<160> 14

<170> PatentIn Ver. 3.3

<210> 1

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1

cggtttccgt gcgttacga

19

<210> 2

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oligonucleotide

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ctggtttcat ctgattactt tcca

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<210> 3

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oligonucleotide

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oligonucleotide

<400> 4
agtcgccta tacgctgct actttcac 28

<210> 5
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oligonucleotide

<400> 5
tgctactttc acgctggaaa gta 23

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oligonucleotide

<400> 6
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<210> 7
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<220>
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oligonucleotide

<400> 7
aggggcatga tgctttcttt ttgttc 26

<210> 8
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<400> 8
 cttgctcggc agagcgagtg ata 23

<210> 9
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 <223> Description of Artificial Sequence: Synthetic
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<400> 9
 gccgagcaag aacaaaaaga aagca 25

<210> 10
 <211> 680
 <212> DNA
 <213> Neisseria gonorrhoeae

<400> 10
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 ttttatattg aaatattatt tttcaaaata aaaatcccaa aatttaccg aaatttggtc 120
 cgaaaaatgg ttttttttcg gggggggtaa ttggagactg attgggtgtt tgcccgatgt 180
 ttttagcaaa tttacaaaag gaagcogata tgcgaaaaaa acttaccgcc ctcgatttgt 240
 ccgcactgcc gtttgcggca gttgccgatg tcggcctgta cggcgaagtc aaagctggtg 300
 tggaaggcag gaacatccgg ctgcagttga ccgagccacc ctcagaaggt caaacgggca 360
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 caaacgccac gacggtatgc 680

<210> 11
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
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<400> 11
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<210> 12
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 oligonucleotide

<400> 12
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<210> 13
 <211> 292
 <212> DNA
 <213> *Neisseria gonorrhoeae*

<400> 13
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 agatgaaacc agttccggct gttgtcggca agccgggggc ggatgtgtat tatgccggtc 180
 tgaattacaa aaatggcggc tttttcggaa attatgccct taaatatgcg aaacacgcca 240
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<210> 14
 <211> 378
 <212> DNA
 <213> *Neisseria gonorrhoeae*

<400> 14
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 aatctcgct tggcggctca gttggatttg tctgaaaatg ccgacaaaac caaaaacagt 120
 acgaccgaaa ttgccgccac tgcttcctac cgcttcggta atacagtccc gcgcatcagc 180
 tatgcccatt gtttcgactt tgcgaacgc agtcagaaac gcgaacatac cagctatgat 240
 caaatcatcg ccggtgtcga ttacgatttt tccaagcgca cttccgccat catgtctgcc 300
 gcttggtgta aacgaaatac cggcatcggc aactacactc aaattaatgc cgcctccggt 360
 ggtctgcgcc acaaattc 378